

Regional immunity in tissue homeostasis and diseases

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The immune system functions in the organ/tissue of the body. The immune cell differentiation and function are influenced by the organ/tissue microenvironments in which they reside, and the interaction of immune cells with the organ/tissue microenvironments may affect and even determine the outcome of the immune responses (Hu and Pasare, 2013; Zajac and Harrington, 2014). It has been increasingly appreciated that the immunologic characterization from professional immune organs including thymus, bone marrow, spleen, and lymph node, is greatly distinguished from that in mucosal immune organs including skin, intestine, lung, pancreas, uterus and kidney, and other organs such as liver, bone, brain, maternal-fetal interface, and so on. Each organ/tissue is characterized with its own anatomy and microenvironment, in which immune cell populations and subsets may have unique functional molecules, forming “unique regional immune features”. The regional immune responses play an important role in tissue homeostasis and are extensively involved in the pathogenesis of the major diseases related to each tissue/organ.

With the research progressed, a more clear description of the regional immune features has been presented, such as

novel tissue-resident immune cell subsets, organ-specialized immune recognition, organ-specific factors instructing regional immune responses in liver, lung, intestine, bone, brain, uterus, adipose, kidney, tumor, etc. For example, specialized tissue-resident macrophages, including Kupffer cells in the liver, alveolar macrophages in the lung, microglia in the brain, osteoclasts in the bone, and histiocytes in the intestinal connective tissue, exhibit distinct functions due to their resident tissue (Murray and Wynn, 2011). Specialized tissue-resident regulatory T cells in the visceral adipose tissue (VAT Tregs), muscle Tregs, bone Tregs and skin memory Tregs, have also been reported with their unique functions in local tissues (Zhou et al., 2015). Innate immune cells continuously receive signals from the resident environment, such as commensal microbiota signals in mucosal organs, which will influence the outcome of immune responses. In this issue, Zhang et al. reviewed the innate recognition of microbial-derived signals in immunity and inflammation (Zhang and Liang, 2016), which will help us understand the local immune responses.

Liver is an unique organ with predominant innate immunity, enriched with Kupffer cells, NK cells, NKT cells and $\gamma\delta$ T cells (Gao et al., 2008). Current studies have revealed that NK cells are not a homogeneous population, but instead consist of distinct subsets including conventional NK (cNK) cells and tissue-resident NK (trNK) cells. In this issue, Peng

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and Tian summarized the phenotypic and functional features of liver trNK cells (Peng and Tian, 2016). In mice, liver NK cells can be divided into two distinct subsets: CD49a⁺DX5⁺ cNK and CD49a⁺DX5⁻ trNK cells which reside in the liver sinusoid blood and confer memory-like responses in contact hypersensitivity (CHS) models (Peng et al., 2013). Liver trNK cells and cNK cells require different transcription factors for their development, thus representing separate developmental lineages (Constantinides et al., 2014; Daussy et al., 2014; Klose et al., 2014; Mackay et al., 2016). They also overviewed recent advances in human liver trNK cells and discussed the striking shared hallmarks of trNK cells in different tissues. Human hepatic CD56^{bright} NK cells express trNK cell markers such as CD69 and CXCR6, and exhibit unique transcriptional profiles compared to cNK cells (Hudspeth et al., 2016). Similar to murine liver trNK cells, a proportion of human hepatic CD56^{bright} NK cells with the expression of CD49a are T-bet⁺Eomes⁻ (Marquardt et al., 2015). Additionally, in this issue Han et al. identified a new subset of inflammation-induced CD69⁺ Kupffer cells during *Listeria monocytogenes* infection (Zhang et al., 2016). Hepatic CD69⁺ Kupffer cells expressed higher levels of CD11b, but produced lower levels of F4/80, which could not be depleted by clodronate liposome administration. Hepatic CD69⁺ Kupffer cells suppressed Ag-nonspecific and OVA-specific CD4⁺ T cell proliferation through membrane-bound TGF- β 1 (mTGF- β 1). CD69⁺ Kupffer cells may contribute to protect host from pathological injury by preventing immune over-activation.

During human pregnancy, an orchestrated evolutionary maternal adaption toward tolerance of the semiallogeneic fetus is required. In the decidua, remodeling of the immune system involves NK cells, macrophages, T cells and DCs to alter the local microenvironment (Fu et al., 2014; Lima et al., 2014). NK cells as the largest population of immune cells during the first trimester in human decidua, accounting for 50%–70% of the total number of immune cells after 9–12 weeks of pregnancy, showed a close reaction to invasive fetal trophoblasts and stromal cells (Colucci and Kieckbusch, 2015; Tao et al., 2015). Evidence shows decidual NK cells act as sentinel cells to control local inflammation and tolerance, and mediate the regulation of fetal growth. In this issue, Fu et al. summarized the concepts and rationale for the remodeled immune system during pregnancy, and the formation and education of decidual NK cells (Fu and Wei, 2016). Most decidual NK cells are CD56^{bright}CD16⁻ NK cells, which express CXCR4, and can be selectively recruited into the decidua by interaction with chemokine CXCL12 secreted by DSCs and EVT (Hanna et al., 2003; Wu et al., 2005).

Adipose tissues, including white, brown and beige fat, play pivotal roles in metabolic homeostasis (Rosen and Spiegelman, 2014). Macrophages, mast cells, neutrophils, eosinophils, ILC2, T cells and

B cells have all been found in adipose tissue and were involved in obesity-related metabolic dysfunctions (Cipolletta et al., 2012; Feuerer et al., 2009; Liu et al., 2009; Lynch et al., 2012; Molofsky et al., 2013; Talukdar et al., 2012; Winer et al., 2009; Winer et al., 2011; Wu et al., 2011). Particularly, macrophages as the critical effector cells in orchestrating inflammation have been most extensively investigated to explore their roles in obesity-associated metabolic inflammation and insulin resistance (McNelis and Olefsky, 2014). In this issue, Qiu et al. reviewed the metabolic activation and functional properties of adipose tissue macrophage (ATM) including M1 and M2 macrophages (Qiu et al., 2016), which would help us better understand the pathogenic and/or protective roles of ATM and the translational opportunities for developing novel therapeutics against obesity, type 2 diabetes and other related metabolic diseases.

Osteoimmunology was firstly introduced in 2000, indicating the bone and immune system have an intense interaction and should be considered as a functional unit (Arron and Choi, 2000). Osteoblasts (OBs) contribute to hematopoietic progenitor growth and HSC pool maintenance through Notch signal and osteopontin (OPN) production (Calvi et al., 2003; Nilsson et al., 2005). The development and response of immune cells associate to bone remodeling; on the other hand, activated immune cells are involved in bone metabolism regulation, and mediate osteoporosis and bone erosion under pathological conditions such as rheumatoid arthritis (RA), spondyloarthropathy (SPA), osteoporosis (OP), and periodontal disease (PD). In this issue, Zhao et al. presented current understanding of osteoimmunology and emphasized on the RANK/RANKL/OPG signaling in bone remodeling and immune responses (Zhao et al., 2016). Recent research progress in the interactions between T cells, B cells, DCs and bone cells, in the role of several kinds of cytokines including M-CSF, TNF- α , IL-1, IFN- γ , IL-4, IL-10, IL-6, TGF- β , OPN and complement system in bone homeostasis were also summarized in this review, particularly their pathogenesis in major diseases.

The central nervous system (CNS) as an immune privileged organ, owns its local tissue barrier and immunosuppressive microenvironment. Interestingly, Glial cells including activated microglia and astrocytes, participate in local immune regulation within the CNS. Further, a specialized CNS-cytokine network was constituted and regulated the development and recovery from autoimmune diseases within the CNS (Xiao and Link, 1998). Additionally, numerous myeloid cell subsets were found in CNS-adjointing tissues, namely the meninges, the perivascular space, and the choroid plexus (Brendecke and Prinz, 2015). These immune cells have been demonstrated to be involved in the initiation, progression and resolution of multiple sclerosis (MS). In this issue, Yan et al. reported a unique group of patients with neu-

romyelitis optica spectrum disorder (NMOSD) who carried autoantibodies of aquaporin-4 (AQP4) and myelin-oligodendrocyte glycoprotein (MOG) (Yan et al., 2016). They found NMOSD patients carrying both anti-AQP4 and anti-MOG antibodies exhibited combined features of prototypic NMO and relapsing-remitting form of MS, whereas NMOSD with only anti-MOG antibody exhibited an “intermediate” phenotype between NMOSD and MS, indicating that anti-MOG antibody might be pathogenic in NMOSD patients and that determination of anti-MOG antibody may be instructive for management of NMOSD patients.

The kidney has its unique organizational structure and physiological functions. It has been known recruitment of leukocytes into the kidney occurs in capillaries of the renal glomeruli and in postcapillary venules of the renal cortex, which is tissue-specific (Rossaint and Zarbock, 2013). Kidney diseases are protean, such as acute kidney injury (AKI), autoimmune glomerulonephritis, and rejection of kidney transplants. Strong evidence demonstrates immune cells, predominantly DCs and macrophages determine both renal tissue injury and subsequent reparative responses (Rogers et al., 2014). However, direct kidney injury mediated by renal infiltrated T cells has not been clearly elucidated in humans. In this issue, Hu et al. detected different types of T cells and macrophages in the renal biopsy tissues from patients of anti-glomerular basement membrane (GBM) disease (Hu et al., 2016). They found that the distribution of T cells was predominant in the peri-glomerular and interstitial areas, particularly, the infiltrating CD8⁺ T cells correlated with the percentage of ruptured Bowman’s capsules, which may participate in the pathogenic mechanism of glomerular damage.

Tumor tissue is also a kind of important regional immune response area. Tumor associated immune cells undergo phenotypic and functional alterations due to the tumor microenvironments (Liu and Cao, 2015a; Liu and Cao, 2015b). In this issue, Xie et al. reported lactic acid in tumor microenvironments inhibited the production of IFN- γ and IL-4 in NKT cells, and these dysfunctions of NKT cells could simply be induced by low extracellular pH through inhibiting mTOR signaling (Xie et al., 2016). It is interesting that tumor acidic microenvironments could regulate immune cell response through modulating cellular metabolisms. Additionally, in the tumor microenvironment a variety of immune factors will alter and aggravate the disease process. Also, Lu et al. reported IL-17 could inhibit the formation of malignant pleural effusion (MPE) via promoting the differentiation of Th9 and Th2 cells in the tumor microenvironments (Lu et al., 2016).

Based on the characteristics of regional immune responses, the future research will focus on discovering novel immune cell subsets and their origin and development in certain types of tissue/organs, exploring the key functional molecules of

the tissue-resident immune cell subsets and their interactions with tissue/organ microenvironments, clarifying their immunological mechanisms in organ-related major diseases, explaining the local environmental factors involved in the tissue residency, which will provide a theoretical basis for the prevention and control of these major diseases and reveal the possible immune intervention targets and immune treatment.

Compliance and ethics The author(s) declare that they have no conflict of interest.

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Biographical Sketch



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Xuetao Cao is professor and President of Chinese Academy of Medical Sciences & Peking Union Medical College. He received Ph.D. from Second Military Medical University (Shanghai, China) in 1990, became Professor in Immunology in 1992 and Director of the Institute of Immunology in 2001 at the same University. He is Professor and Director of National Key Laboratory of Medical Immunology (2006.5–). He was elected as member of Chinese Academy of Engineering in 2005, Foreign Associate of German Academy of Sciences in 2013, and Foreign Member of EMBO in 2015. His major interests are innate response, inflammation and tumor immunotherapy. As corresponding author, he published more than 230 original papers in peer-reviewed journals including *Cell*, *Nature*, *Science*. He is Co-Editor-in-Chief of *Cellular and Molecular Immunology*, editorial board member of *Cell*, *Annual Reviews of Immunology*, *Science Translational Medicine*, *eLife*, *Cell Research*, etc.



Yongyan Chen is an associate Professor of University of Science & Technology of China (USTC). As a member of Zhigang Tian's lab in USTC, she has been attempting to study the innate immunity of liver, including the biology of the overwhelming innate immune cells NK cells and NKT cells in the liver, and their roles during HBV infection and HBV-associated liver diseases; and the mechanisms of HBV-induced liver immunotolerance.



Qunyan Lyu, professor and Division Director, Division IV of Department of Health Sciences, National Natural Science Foundation of China. She received her Ph.D. in Microbiology in 1993, and has been responsible for the management of natural science fund in the field of immunology or medical immunology for about 20 years. She has engaged herself in implementing the funding plans to support basic research, identifying and fostering scientific talents, accepting project applications, organizing peer review process, administering funded projects etc, in the field of immunology or medical immunology. As a professional scientific manager, she is also playing active roles in formulation of the priority funding areas, funding schemes, major research plans, annual funding arrangements as well as other disciplinary development strategies for the field of immunology or medical immunology.